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OM protein - protein search, using sw model

Run on: February 11, 2003, 19:45:09 ; Search time 13.1143 Seconds
(without alignments)
991.661 Million cell updates/sec

Title: US-09-497-967-6
Perfect score: 2342
Sequence: 1 MYNILLIILISLFINELRA.....STTRAKFLISILLFISFYLL 442

Scoring table: BLOSUM62

Gapop 10.0 , Gapext 0.5

Searched: 262574 seqs, 29422922 residues

Total number of hits satisfying chosen parameters: 262574

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database :

- 1: /cgn2_6/ptodata/1/1aa/5A.COMB.pep.*
- 2: /cgn2_6/ptodata/1/1aa/5B.COMB.pep.*
- 3: /cgn2_6/ptodata/1/1aa/6A.COMB.pep.*
- 4: /cgn2_6/ptodata/1/1aa/6B.COMB.pep.*
- 5: /cgn2_6/ptodata/1/1aa/PCTUS.COMB.pep.*
- 6: /cgn2_6/ptodata/1/1aa/backfiles1.pep.*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Query Match	Score	Length	ID	Description
1	234	10.0	1652	4	US-09-627-650B-1
2	234	10.0	1652	4	US-09-436-063C-1
3	233.5	10.0	2508	4	US-09-627-650B-7
4	233.5	10.0	2508	4	US-09-436-063C-7
5	233.5	10.0	2544	4	US-09-627-650B-3
6	233.5	10.0	2544	4	US-09-436-063C-3
7	233.5	10.0	2601	4	US-09-627-650B-9
8	233.5	10.0	2601	4	US-09-436-063C-9
9	230	9.8	1917	4	US-09-627-650B-5
10	230	9.8	1917	4	US-09-436-063C-5
11	219.5	9.4	1128	4	US-09-627-650B-11
12	219.5	9.4	1128	4	US-09-436-063C-11
13	219	9.4	341	2	US-08-209-521-11
14	202.5	8.6	2088	4	US-09-548-372D-13
15	202.5	8.6	2088	4	US-09-548-367D-13
16	195.5	8.3	801	1	US-07-906-349A-6
17	194.5	8.1	1400	4	US-08-630-915A-37
18	190.5	8.1	3111	2	US-08-460-309-4
19	190.5	8.1	3111	2	US-08-125-077-4
20	186	7.9	3075	2	US-08-460-309-5
21	186	7.9	3075	2	US-08-125-077-5
22	177.5	7.6	2211	4	US-09-738-884-1
23	173.5	7.4	288	2	US-08-525-940-15
24	173.5	7.4	288	2	US-08-976-838-15
25	172	7.3	799	2	US-08-525-940-23
26	172	7.3	799	2	US-08-976-838-23
27	172	7.3	881	2	US-08-525-940-21

28	172	7.3	881	2	US-08-976-838-21	Sequence 21, Appl
29	172	7.3	915	2	US-08-525-940-18	Sequence 18, Appl
30	172	7.3	915	2	US-08-976-838-18	Sequence 18, Appl
31	172	7.3	915	4	US-09-214-555B-2	Sequence 2, Appl
32	172	7.3	915	4	US-09-214-555B-7	Sequence 7, Appl
33	168.5	7.2	288	1	US-08-368-852-15	Sequence 15, Appl
34	162.5	6.9	1345	2	US-08-977-767-3	Sequence 3, Appl
35	154.5	6.6	2556	3	US-08-083-590A-20	Sequence 20, Appl
36	154.5	6.6	2556	3	US-08-532-384-20	Sequence 20, Appl
37	153.5	6.6	1417	4	US-08-900-230-3	Sequence 3, Appl
38	151.5	6.5	3224	2	US-08-705-660-34	Sequence 34, Appl
39	151.5	6.5	3224	3	US-08-989-045-34	Sequence 34, Appl
40	150.5	6.4	1111	1	US-08-317-450B-15	Sequence 15, Appl
41	150.5	6.4	1111	4	US-08-800-593-15	Sequence 15, Appl
42	150.5	6.4	1193	1	US-08-317-450B-13	Sequence 13, Appl
43	150.5	6.4	1193	4	US-08-800-593-13	Sequence 13, Appl
44	149.5	6.4	2556	1	US-08-185-432-17	Sequence 17, Appl
45	149.5	6.4	2556	4	US-08-899-232-2	Sequence 2, Appl

ALIGNMENTS

RESULT 1
US-09-627-650B-1
; Sequence 1, Application US/09627650B
; Patent No. 6406872
; GENERAL INFORMATION:
; APPLICANT: Bamber, Bruce
; TITLE OF INVENTION: Nematode Neuromuscular Junction GABA Receptors and
; TITLE OF INVENTION: Methods Related Thereto
; FILE REFERENCE: 21101.000903
; CURRENT APPLICATION NUMBER: US/09/627,650B
; CURRENT FILING DATE: 2000-07-28
; PRIOR APPLICATION NUMBER: 09/436,063
; PRIOR FILING DATE: 1999-11-08
; PRIOR APPLICATION NUMBER: 60/107,727
; PRIOR FILING DATE: 1998-11-09
; NUMBER OF SEQ ID NOS: 50
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 1
; LENGTH: 1652
; TYPE: PRT
; ORGANISM: Caenorhabditis elegans
US-09-627-650B-1

Query Match	10.0%	Score 234;	DB 4;	Length 1652;
Best Local Similarity	26.4%	Pred. No. 2.4e-11;		
Matches 115;	Conservative 14;	Mismatches 211;	Indels 96;	Gaps 17;
QY	20	AVPCPDGTQTQAGLTDVGAADLGTVCVNCR-----PNFYNGGAAGCAAGTGTCCAAATGCAAGCTGAAGCTGTTCCTCAA	74	
Db	580	ATACTAGTCAAGATTAAACAGTCACTGCAACGTGTCCAAATGCAAGCTGAAGCTGTTCCTCAA	639	
QY	75	RG-----ICVPCQINRVGSVTNAGDLAT--LATQC-STOCPTGTALDDGVTDVDFDRSAQ	126	
Db	640	TGGACTCTCAACACTGTAAGTGAAGTATGCG--GTATTCGACGCCGGAAT	697	
QY	127	CVKCKPNFYNGSGPGEAPGVQVFAAGAAAGVAAVT-SQCVPCLNKNKNDSPATAG----	182	
Db	698	CGAGTACAAATGCT--GTACGTGGAAGGAGCGCAATTTGTCGAC--AGCGGTCAAGGCCGA	754	
QY	183	-----AQAANLATOCSNOCPTGTVLDDGVTLVNT-SATLCVKCRPNFYNGSGPGEAPG	236	
Db	755	CGGCAACATCGAAGCTGCGAGTTATAAATTCATAAAATCTGC-----	797	
QY	237	VQVFAAGAAAGVAAVTSQVPCQINKNKNDSPATAQAANLATQCSQCTGCTATODGVTL	296	
Db	798	---CAAAAACGGACACTTGC--CAGCACTTCATCGGGACCTACTCTCGTCTACGGGTTA	852	
QY	297	VFSNSTQ-----CSQCIANYFFNFGKSKQCLKCPVSKTTPAHAFNPATQATQ	348	

Db 853 GTTTCATATTGATCGGAC-----AGCGGCTCTTACTTCTTCAAATATTTT 901

QY 349 CLTTCPAGTVLDDGTSTNFVASTECTKCSAGFFASKTTGTFAGTDTCTCTCKLTSAT 408

Db 902 C---C-----CTGCCAGCTCGTCG-TAGTTTATCA---TGGAT 934

QY 409 AKVYAEATQKVOCAST 424

Db 935 CTCATTCTGGATCAAT 950

RESULT 2

US-09-436-063C-1

; Sequence 1, Application US/09436063C

; Patent No. 6407210

; GENERAL INFORMATION:

; APPLICANT: Bamber, Bruce

; APPLICANT: Jorgensen, Erik

; TITLE OF INVENTION: Nematode Neuromuscular Junction GABA Receptors and

; TITLE OF INVENTION: Methods Related Thereto

; FILE REFERENCE: P-1095corrected

; CURRENT APPLICATION NUMBER: US/09/436,063C

; CURRENT FILING DATE: 1999-11-08

; PRIOR APPLICATION NUMBER: 60/107727

; PRIOR FILING DATE: 1998-11-09

; NUMBER OF SEQ ID NOS: 18

; SOFTWARE: PatentIn Ver. 2.1

; SEQ ID NO 1

; LENGTH: 1652

; TYPE: PRT

; ORGANISM: Caenorhabditis elegans

US-09-436-063C-1

Query Match 10.0%; Score 234; DB 4; Length 1652;

Best Local Similarity 26.4%; Pred.No. 2.4e-11;

Matches 115; Conservative 14; Mismatches 211; Indels 96; Gaps 17;

QY 20 AVPCPDGTGTQAGLTDVGAADLGLTCVNCR-----PNFYINGGAAQGEANGNOPFAANNA 74

Db 580 ATACTAGTCAAGATTACAGTCACTGCAAGCTGTCCAATGGACCTGAAGCTGTTCCCAA 639

QY 75 RG-----ICVPCQINRVGSVFNAGDLAT--LATOC-STOCPTGTALDDGVTDFEDRSAQ 126

Db 640 TGGACTCTCAACACTGTAACACTGGAAATTTGAAAGCTATGC--GTATTCGACGCCGCAAT 697

QY 127 CVKCKPNFYNGSGPOGAPGVQVFAAGAAAGVAAYTSQVPCQLNKNDSPATAG----182

Db 698 CGAGTACAATGTG--GTACGTGCGAAGGCGGAATTTGCGAC-AGCGGTCAAGGCCGA 754

QY 183 -----AQAANIATQCSNOCTPGTVLDDGTVLFWNT-SATLCVKCKPNFYNGSGPOGEAFG 236

Db 755 CGCGAACATCGAACTGTCGATATATAAATTCATAAATCTGC-----797

QY 237 VQVFAAGAAAAGVAAYTSQVPCQINKNDSPATAGAAANLATQCTOCPTGTATQDGVTL 296

Db 798 ---CAAAAACGGACACTTGC--CAGCACTTCATCGGGACCTACTCTCGTCTACGGGTTA 852

QY 297 VFSNSSTQ-----CSOCIANYFPNGNFEAGKSQCLKCPVSKYTTFAHPAGNTATQATQ 348

Db 853 GTTTCATATTGATCGGAC-----AGCGGCTTCTACTTCTTCAAATATTTT 901

QY 349 CLTTCPAGTVLDDGTSTNFVASTECTKCSAGFFASKTTGTFAGTDTCTCTCKLTSAT 408

Db 902 C---C-----CTGCCAGCTCGTCG-TAGTTTATCA---TGGAT 934

QY 409 AKVYAEATQKVOCAST 424

Db 935 CTCATTCTGGATCAAT 950

RESULT 3

US-09-627-650B-7

; Sequence 7, Application US/09627650B

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; ORGANISM: Caenorhabditis elegans
US-09-436-063C-7

Query Match          10.0%; Score 233.5; DB 4; Length 2508;
Best Local Similarity 26.7%; Pred. No. 4.5e-11;
Matches 112; Conservative 13; Mismatches 188; Indels 107; Gaps 17;

QY 27 TQTQAGLTDVGAADLGTVCNCRPNFYNGGAAQGEANGNPPFAANNAARGICVPCQINRV 86
Db 1500 TCTAAGCTTAAACTCGTC-----GGACAAGAT-----CCAGGAGAACGACAAGTG 1544

QY 87 GSVTNAGDLATATQC-----STQCPTGTALDDGVTDVDRSAACVKCKPNFYNGGSPQ 142
Db 1545 GCAGCAG-----ATCCACTGATGC--GTATTTCGACGCCGGAATCGAGTACAAATGGT-- 1595

QY 143 GEAPGVQVFAAGAAAGVAATVSOQVPCQLNKNDSPATAG-----AQNATQCSN 193
Db 1596 GTACGTGCAAGGAGCCGAATTTGTCGAC-AGCGGTCAAGGCCGACGCGAATCGAAGTG 1654

QY 194 QCPTGTDLDDGVTLVFNVT-SATLCVKCRPNFYNGGSPQGEAPGVQVFAAGAAAGVAAV 252
Db 1655 TCGAGTTATAAATTCATAAATCTGC-----C-----CAAAAACGGACAC 1694

QY 253 TSOCVPCQINKNDSPATAGAAANLATQCSTQCPTGTATQDGTATQDGTATQDGTATQDGT 304
Db 1695 TTGC--CAGCAGCTTACATCGGGGACCTACTCTGCTCTACGGGTTAGTTTCATATTTGATCG 1752

QY 305 CSOCIANYFFNGFNFEAKGSOCLKCPVSKTTPAHAPGNATATQATQCLTTCAGTVLDDGTS 364
Db 1753 CGAC-----AGCGGCTTCTACTTCTTCAAAATATTTTC--C-----1787

QY 365 TNFVASATECTKCSAGFFASKTTGTTAGTDTCTECTKLTSGATAKVYAEATQKVOCAST 424
Db 1788 -----CTGCCAGCCTCGTCG-TAGTTTATCA---TGGATCTCATCTCGGATCAAT 1834

RESULT 5
US-09-627-650B-3
; Sequence 3, Application US/09627650B
; Patent No. 6406872
; GENERAL INFORMATION:
; APPLICANT: Bamber, Bruce
; APPLICANT: Jorgensen, Erik
; TITLE OF INVENTION: Nematode Neuromuscular Junction GABA Receptors and
; FILE OF INVENTION: Methods Related Thereto
; FILE REFERENCE: 21101.000903
; CURRENT APPLICATION NUMBER: US/09/627,650B
; PRIOR FILING DATE: 2000-07-28
; PRIOR APPLICATION NUMBER: 09/436,063
; PRIOR FILING DATE: 1999-11-08
; PRIOR APPLICATION NUMBER: 60/107,727
; PRIOR FILING DATE: 1998-11-09
; NUMBER OF SEQ ID NOS: 50
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 3
; LENGTH: 2544
; TYPE: PRT
; ORGANISM: Caenorhabditis elegans
US-09-627-650B-3

Query Match          10.0%; Score 233.5; DB 4; Length 2544;
Best Local Similarity 26.7%; Pred. No. 4.6e-11;
Matches 112; Conservative 13; Mismatches 188; Indels 107; Gaps 17;

QY 27 TQTQAGLTDVGAADLGTVCNCRPNFYNGGAAQGEANGNPPFAANNAARGICVPCQINRV 86
Db 1536 TCTAAGCTTAAACTCGTC-----GGACAAGAT-----CCAGGAGAACGACAAGTG 1580

QY 87 GSVTNAGDLATATQC-----STQCPTGTALDDGVTDVDRSAACVKCKPNFYNGGSPQ 142
Db 1581 GCAGCAG-----ATCCACTGATGC--GTATTTCGACGCCGGAATCGAGTACAAATGGT-- 1631

QY 143 GEAPGVQVFAAGAAAGVAATVSOQVPCQLNKNDSPATAG-----AQNATQCSN 193
Db 1632 GTACGTGCAAGGAGCCGAATTTGTCGAC-AGCGGTCAAGGCCGACGCGAATCGAAGTG 1690

QY 194 QCPTGTDLDDGVTLVFNVT-SATLCVKCRPNFYNGGSPQGEAPGVQVFAAGAAAGVAAV 252
Db 1691 TCGAGTTATAAATTCATAAATCTGC-----C-----CAAAAACGGACAC 1730

QY 253 TSOCVPCQINKNDSPATAGAAANLATQCSTQCPTGTATQDGTATQDGTATQDGTATQDGT 304
Db 1731 TTGC--CAGCAGCTTACATCGGGGACCTACTCTGCTCTACGGGTTAGTTTCATATTTGATCG 1788

QY 305 CSOCIANYFFNGFNFEAKGSOCLKCPVSKTTPAHAPGNATATQATQCLTTCAGTVLDDGTS 364
Db 1789 CGAC-----AGCGGCTTCTACTTCTTCAAAATATTTTC--C-----1823

QY 365 TNFVASATECTKCSAGFFASKTTGTTAGTDTCTECTKLTSGATAKVYAEATQKVOCAST 424
Db 1824 -----CTGCCAGCCTCGTCG-TAGTTTATCA---TGGATCTCATCTCGGATCAAT 1870

; ORGANISM: Caenorhabditis elegans
US-09-436-063C-3
; Sequence 3, Application US/09436063C
; Patent No. 6407210
; GENERAL INFORMATION:
; APPLICANT: Bamber, Bruce
; APPLICANT: Jorgensen, Erik
; TITLE OF INVENTION: Nematode Neuromuscular Junction GABA Receptors and
; FILE OF INVENTION: Methods Related Thereto
; FILE REFERENCE: P-1095corrected
; CURRENT APPLICATION NUMBER: US/09/436,063C
; PRIOR FILING DATE: 1999-11-08
; PRIOR APPLICATION NUMBER: 60/107727
; PRIOR FILING DATE: 1998-11-09
; NUMBER OF SEQ ID NOS: 18
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 3
; LENGTH: 2544
; TYPE: PRT
; ORGANISM: Caenorhabditis elegans
US-09-436-063C-3

Query Match          10.0%; Score 233.5; DB 4; Length 2544;
Best Local Similarity 26.7%; Pred. No. 4.6e-11;
Matches 112; Conservative 13; Mismatches 188; Indels 107; Gaps 17;

QY 27 TQTQAGLTDVGAADLGTVCNCRPNFYNGGAAQGEANGNPPFAANNAARGICVPCQINRV 86
Db 1536 TCTAAGCTTAAACTCGTC-----GGACAAGAT-----CCAGGAGAACGACAAGTG 1580

QY 87 GSVTNAGDLATATQC-----STQCPTGTALDDGVTDVDRSAACVKCKPNFYNGGSPQ 142
Db 1581 GCAGCAG-----ATCCACTGATGC--GTATTTCGACGCCGGAATCGAGTACAAATGGT-- 1631

QY 143 GEAPGVQVFAAGAAAGVAATVSOQVPCQLNKNDSPATAG-----AQNATQCSN 193
Db 1632 GTACGTGCAAGGAGCCGAATTTGTCGAC-AGCGGTCAAGGCCGACGCGAATCGAAGTG 1690

QY 194 QCPTGTDLDDGVTLVFNVT-SATLCVKCRPNFYNGGSPQGEAPGVQVFAAGAAAGVAAV 252
Db 1691 TCGAGTTATAAATTCATAAATCTGC-----C-----CAAAAACGGACAC 1730

QY 253 TSOCVPCQINKNDSPATAGAAANLATQCSTQCPTGTATQDGTATQDGTATQDGTATQDGT 304
Db 1731 TTGC--CAGCAGCTTACATCGGGGACCTACTCTGCTCTACGGGTTAGTTTCATATTTGATCG 1788

QY 305 CSOCIANYFFNGFNFEAKGSOCLKCPVSKTTPAHAPGNATATQATQCLTTCAGTVLDDGTS 364
Db 1789 CGAC-----AGCGGCTTCTACTTCTTCAAAATATTTTC--C-----1823

QY 365 TNFVASATECTKCSAGFFASKTTGTTAGTDTCTECTKLTSGATAKVYAEATQKVOCAST 424
Db 1824 -----CTGCCAGCCTCGTCG-TAGTTTATCA---TGGATCTCATCTCGGATCAAT 1870
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; NUMBER OF SEQ ID NOS: 18
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 9
; LENGTH: 2601
; TYPE: PRT
; ORGANISM: Caenorhabditis elegans
US-09-436-063C-9

Query Match          10.0%; Score 233.5; DB 4; Length 2601;
Best Local Similarity 26.7%; Pred. No. 4.7e-11;
Matches 112; Conservative 13; Mismatches 188; Indels 107; Gaps 17;

QY 27 TOTOAGLTDVGAADLGTVCNCRPNFYNGAAGGANGNQPFAANNAARGICVPCQINRV 86
    ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 1593 TCTAAGCTTAAACTCGTC-----GGACAAGAT-----CCAGGAGAAGCAAGTG 1637
    ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||

QY 87 GSVTNAGDLATATQC-----STQCPTGTALDGVTDVDFRSAAQCCKPNFYNGGSPQ 142
    ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 1638 GCAGCAG-----ATCCACTGATGC--GTATTGACGCGCGGAATCGAGTACAAATGGT-- 1688
    ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||

QY 143 GEAPGVQVFAAGAAAAGVAAVTSQVPCOLNKNDSPATAG-----AQNLATQCSN 193
    ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 1689 GTAGTCGAAGGAGCGCAATGTTGCGAC-AGCGGTCAAGCGCGCAACATCGAACTG 1747
    ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||

QY 194 QCPTGTVLDGVTLVFN--SATLCVKCRPNFYNGSPGEGAPGVQVFAAGAAAAGVAAV 252
    ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 1748 TCGAGTTATAAATTCATAAAATCTGC-----CAAAAACGCGAC 1787
    ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||

QY 253 TSQVPCQINKNDSPATAGAAANLATQCSQCPTGTAIODGVTLVFSNSTQ----- 304
    ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 1788 TTGC--CAGCACTTCATCGGGACCTACTCTGCTACGGGTAGTTTCATATTGATCG 1845
    ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||

QY 305 CSOCIANYFFNGNFEAGKSQCLKCPVSKTTPAHAPGNATQATQCLTTCPAGTVLDGTS 364
    ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 1846 CGAC-----ACGGCCTTCTACTTCTTCAAATATTTTC--C----- 1880
    ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||

QY 365 TNFVASATECTKCSAGFFASKTTGFTAGTDTCTECTKLTSGATAKYAEATQKVOCAST 424
    ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 1881 -----CTGCCAGCCTCGTCG--TAGTTTATCA---TGGATCTCATCTCGATCAAT 1927
    ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||

RESULT 9
US-09-627-650B-5
; Sequence 5, Application US/09627650B
; Patent No. 6406872
; GENERAL INFORMATION:
; APPLICANT: Bamber, Bruce
; APPLICANT: Jorgensen, Erik
; TITLE OF INVENTION: Nematode Neuromuscular Junction GABA Receptors and
; TITLE OF INVENTION: Methods Related Thereto
; FILE REFERENCE: 21101.0009U3
; CURRENT APPLICATION NUMBER: US/09/627,650B
; CURRENT FILING DATE: 2000-07-28
; PRIOR APPLICATION NUMBER: 09/436,063
; PRIOR FILING DATE: 1999-11-08
; PRIOR APPLICATION NUMBER: 60/107,727
; PRIOR FILING DATE: 1998-11-09
; NUMBER OF SEQ ID NOS: 50
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 5
; LENGTH: 1917
; TYPE: PRT
; ORGANISM: Caenorhabditis elegans
US-09-627-650B-5

Query Match          9.8%; Score 230; DB 4; Length 1917;
Best Local Similarity 26.1%; Pred. No. 6.4e-11;
Matches 109; Conservative 14; Mismatches 203; Indels 92; Gaps 18;

QY 26 GTQTQAGLTDVGAADLGTVCNCRPNFYNGAAGGANGNQPFAANNAARGICVPC 81
    ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 476 GTTGAATTACGATACACGTTTCTTCAATCTCTCGAGTTTCTAGAGTTGATGATGATTC 535
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; NUMBER OF SEQ ID NOS: 18
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 9
; LENGTH: 2601
; TYPE: PRT
; ORGANISM: Caenorhabditis elegans
US-09-436-063C-9

Query Match          10.0%; Score 233.5; DB 4; Length 2601;
Best Local Similarity 26.7%; Pred. No. 4.7e-11;
Matches 112; Conservative 13; Mismatches 188; Indels 107; Gaps 17;

QY 27 TOTOAGLTDVGAADLGTVCNCRPNFYNGAAGGANGNQPFAANNAARGICVPCQINRV 86
    ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 1593 TCTAAGCTTAAACTCGTC-----GGACAAGAT-----CCAGGAGAAGCAAGTG 1637
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QY 87 GSVTNAGDLATATQC-----STQCPTGTALDGVTDVDFRSAAQCCKPNFYNGGSPQ 142
    ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 1638 GCAGCAG-----ATCCACTGATGC--GTATTGACGCGCGGAATCGAGTACAAATGGT-- 1688
    ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||

QY 143 GEAPGVQVFAAGAAAAGVAAVTSQVPCOLNKNDSPATAG-----AQNLATQCSN 193
    ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 1689 GTAGTCGAAGGAGCGCAATGTTGCGAC-AGCGGTCAAGCGCGCAACATCGAACTG 1747
    ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||

QY 194 QCPTGTVLDGVTLVFN--SATLCVKCRPNFYNGSPGEGAPGVQVFAAGAAAAGVAAV 252
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Db 1748 TCGAGTTATAAATTCATAAAATCTGC-----CAAAAACGCGAC 1787
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QY 253 TSQVPCQINKNDSPATAGAAANLATQCSQCPTGTAIODGVTLVFSNSTQ----- 304
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Db 1788 TTGC--CAGCACTTCATCGGGACCTACTCTGCTACGGGTAGTTTCATATTGATCG 1845
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QY 305 CSOCIANYFFNGNFEAGKSQCLKCPVSKTTPAHAPGNATQATQCLTTCPAGTVLDGTS 364
    ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||
Db 1846 CGAC-----ACGGCCTTCTACTTCTTCAAATATTTTC--C----- 1880
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QY 365 TNFVASATECTKCSAGFFASKTTGFTAGTDTCTECTKLTSGATAKYAEATQKVOCAST 424
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Db 1881 -----CTGCCAGCCTCGTCG--TAGTTTATCA---TGGATCTCATCTCGATCAAT 1927
    ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| ||| |||

RESULT 8
US-09-436-063C-9
; Sequence 9, Application US/09436063C
; Patent No. 6407210
; GENERAL INFORMATION:
; APPLICANT: Bamber, Bruce
; APPLICANT: Jorgensen, Erik
; TITLE OF INVENTION: Nematode Neuromuscular Junction GABA Receptors and
; TITLE OF INVENTION: Methods related Thereto
; FILE REFERENCE: P-1095corrected
; CURRENT APPLICATION NUMBER: US/09/436,063C
; CURRENT FILING DATE: 1999-11-08
; PRIOR APPLICATION NUMBER: 60/107727
; PRIOR FILING DATE: 1998-11-09

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, RESULT 14
, US-09-548-372D-13
, Sequence 13, Application US/09548372D
, Patent No. 6420534
, GENERAL INFORMATION:
, APPLICANT: GURNEY ET AL.
, TITLE OF INVENTION: ALZHEIMER'S DISEASE
, TITLE OF INVENTION: THEREOF
, FILE REFERENCE: 23915/62801
, CURRENT APPLICATION NUMBER: US/09/548
, CURRENT FILING DATE: 2000-04-12

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; PRIOR APPLICATION NUMBER: US 60/155,493
; PRIOR FILING DATE: 1999-09-23
; PRIOR APPLICATION NUMBER: US 09/404,133
; PRIOR FILING DATE: 1999-09-23
; PRIOR APPLICATION NUMBER: PCT/US99/20881
; PRIOR FILING DATE: 1999-09-23
; PRIOR APPLICATION NUMBER: US 60/101,594
; PRIOR FILING DATE: 1998-09-24
; NUMBER OF SEQ ID NOS: 73
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 13
; LENGTH: 2088
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-548-372D-13

Query Match 8.6%; Score 202.5; DB 4; Length 2088;
Best Local Similarity 23.5%; Pred. No. 1.5e-08;
Matches 115; Conservative 16; Mismatches 226; Indels 133; Gaps 19;
QY 23 CPDGTQTQAGLTDVG-----AADLGTVCNCRPNFYNGGAAOGEA-----NGNQPPA 69
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QY 70 ANNAARICVPCQINRVSGVTNAGDLATLQCTGTCPTGTALDDGVTDVDFDRSAAQCVK 129
Db 196 AAGGAAGGCATCCTGCAGTATTGCCAAGAAGTCTACCCCTGAAC-----GCAGATCAC 248
QY 130 CKPNFYNGSGPOGAPGVQVFAAGAAAGVAAVTSQCVPQINKNDSPATAGAANLAT 189
Db 249 C-----AATGTGGTAGAAGCCAAACCAACGAGTGACCATC-----CAGAACTGGT 292
QY 190 QCSNQCPGT-----PTGTALQDGVTLVFNSTQCS-----QCIAN 311
Db 293 GCAAGCGGGCCCAAGCAGTGCAGACCCATCCCACTTTGTGATTCCTACCGCTGC- 351
QY 221 PNFYNGSGPOGAPGVQVFAAGAAAGVAAVTSQC-VPCQINKNDSPATAGAANLATQ 279
Db 352 TTAGTTGGTGAGTTTGT-----AAGTGATGCCCTTCTCGTTC-----CTGACAAGTGCAAAAT 404
QY 280 CSTQC-----PTGTALQDGVTLVFNSTQCS-----QCIAN 311
Db 405 CTTACACGAGGAGAGATGGATGTTTGGAAACTCATCTTCACTGGCACACCGTGCCTG- 464
QY 312 YFEN-----GNFEAGKSQCLKCPVSKTTPA-HAPGNATQATQCLTTCGAGTIVL 359
Db 465 AGAGACATGCAGTGAGAGAGTACCAACTTGCGATGACTACGCGCATGTTGCTGCCTG--C 522
QY 360 DGGTSTNFVASATECTKCSAGFFASKTTGTTAGTDT-----CTECTKLTSGATAKYAE 414
Db 523 GGAATTGACAAGTTC--CGAGGGGTAGAGTTTGTGTGTGTTGCCCTGCTGCTGAAGAAAGTG 580
QY 415 ATQKVOCAST 424
Db 581 A-----CAAT 585

RESULT 15

US-09-548-367D-13
; Sequence 13, Application US/09548367D
; Patent No. 6440698
; GENERAL INFORMATION:
; APPLICANT: GURNEY ET AL.
; TITLE OF INVENTION: ALZHEIMER'S DISEASE SECRETASE, APP SUBSTRATES THEREFOR AND USES
; TITLE OF INVENTION: THEREOF
; FILE REFERENCE: 29915/6280H
; CURRENT APPLICATION NUMBER: US/09/548,367D
; CURRENT FILING DATE: 2000-04-12
; PRIOR APPLICATION NUMBER: US 60/155,493
; PRIOR FILING DATE: 1999-09-23
; PRIOR APPLICATION NUMBER: US 09/404,133
; PRIOR FILING DATE: 1999-09-23
; PRIOR APPLICATION NUMBER: PCT/US99/20881

; PRIOR FILING DATE: 1999-09-23
; PRIOR APPLICATION NUMBER: US 60/101,594
; PRIOR FILING DATE: 1998-09-24
; NUMBER OF SEQ ID NOS: 73
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 13
; LENGTH: 2088
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-548-367D-13

Query Match 8.6%; Score 202.5; DB 4; Length 2088;
Best Local Similarity 23.5%; Pred. No. 1.5e-08;
Matches 115; Conservative 16; Mismatches 226; Indels 133; Gaps 19;
QY 23 CPDGTQTQAGLTDVG-----AADLGTVCNCRPNFYNGGAAOGEA-----NGNQPPA 69
Db 141 CCAGAATGGGAAGTGGGATTCAGATCCATCA-----GGACCAAAACCTGCATTGATACC 195
QY 70 ANNAARICVPCQINRVSGVTNAGDLATLQCTGTCPTGTALDDGVTDVDFDRSAAQCVK 129
Db 196 AAGGAAGGCATCCTGCAGTATTGCCAAGAAGTCTACCCCTGAAC-----GCAGATCAC 248
QY 130 CKPNFYNGSGPOGAPGVQVFAAGAAAGVAAVTSQCVPQINKNDSPATAGAANLAT 189
Db 249 C-----AATGTGGTAGAAGCCAAACCAACGAGTGACCATC-----CAGAACTGGT 292
QY 190 QCSNQCPGT-----PTGTALQDGVTLVFNSTQCS-----QCIAN 311
Db 293 GCAAGCGGGCCCAAGCAGTGCAGACCCATCCCACTTTGTGATTCCTACCGCTGC- 351
QY 221 PNFYNGSGPOGAPGVQVFAAGAAAGVAAVTSQC-VPCQINKNDSPATAGAANLATQ 279
Db 352 TTAGTTGGTGAGTTTGT-----AAGTGATGCCCTTCTCGTTC-----CTGACAAGTGCAAAAT 404
QY 280 CSTQC-----PTGTALQDGVTLVFNSTQCS-----QCIAN 311
Db 405 CTTACACGAGGAGAGATGGATGTTTGGAAACTCATCTTCACTGGCACACCGTGCCTG- 464
QY 312 YFEN-----GNFEAGKSQCLKCPVSKTTPA-HAPGNATQATQCLTTCGAGTIVL 359
Db 465 AGAGACATGCAGTGAGAGAGTACCAACTTGCGATGACTACGCGCATGTTGCTGCCTG--C 522
QY 360 DGGTSTNFVASATECTKCSAGFFASKTTGTTAGTDT-----CTECTKLTSGATAKYAE 414
Db 523 GGAATTGACAAGTTC--CGAGGGGTAGAGTTTGTGTGTGTTGCCCTGCTGCTGAAGAAAGTG 580
QY 415 ATQKVOCAST 424
Db 581 A-----CAAT 585

Search completed: February 11, 2003, 19:49:12
Job time : 24.1143 secs

